

Jadwiga Åliwka

List of Publications by Year in descending order

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Version: 2024-02-01

50
papers

1,691
citations

304743

22
h-index

302126

39
g-index

52
all docs

52
docs citations

52
times ranked

1595
citing authors

#	ARTICLE	IF	CITATIONS
1	Tuber Flesh Colour, Enzymatic Discolouration, Dormancy and Late Blight Resistance of 29 Tuber-Bearing Accessions of <i>Solanum</i> spp.. <i>Potato Research</i> , 2023, 66, 1-21.	2.7	1
2	Quantitative Trait Loci for Resistance to Potato Dry Rot Caused by <i>Fusarium sambucinum</i> . <i>Agronomy</i> , 2022, 12, 203.	3.0	1
3	Late blight resistance genes in potato breeding. <i>Planta</i> , 2022, 255, 127.	3.2	28
4	Population Structure of <i>Phytophthora infestans</i> from a Single Location in Poland Over a Long Period of Time in Context of Weather Conditions. <i>Microbial Ecology</i> , 2021, 81, 746-757.	2.8	10
5	Identification and pathogenicity of <i>Fusarium</i> spp. associated with tuber dry rot and wilt of potato in Algeria. <i>European Journal of Plant Pathology</i> , 2021, 159, 495-509.	1.7	25
6	Analysis of Cytosine Methylation in Genomic DNA of <i>Solanum michoacanum</i> (+) <i>S. tuberosum</i> Somatic Hybrids. <i>Agronomy</i> , 2021, 11, 845.	3.0	6
7	QTLs for potato tuber resistance to <i>Dickeya solani</i> are located on chromosomes II and IV. <i>Plant Pathology</i> , 2021, 70, 1745-1756.	2.4	9
8	Quantitative trait loci for starch-corrected chip color after harvest, cold storage and after reconditioning mapped in diploid potato. <i>Molecular Genetics and Genomics</i> , 2020, 295, 209-219.	2.1	6
9	eQTL mapping of the 12S globulin cruciferin gene PGCRURSE5 as a novel candidate associated with starch content in potato tubers. <i>Scientific Reports</i> , 2020, 10, 17168.	3.3	4
10	Marker-assisted pyramiding of potato late blight resistance genes <i>Rpi-rzc1</i> and <i>Rpi-phu1</i> on di- and tetraploid levels. <i>Molecular Breeding</i> , 2020, 40, 1.	2.1	18
11	Quantitative trait loci analysis of potato tuber greening. <i>Molecular Biology Reports</i> , 2020, 47, 1713-1722.	2.3	8
12	Cytoplasmic diversity of potato relatives preserved at Plant Breeding and Acclimatization Institute in Poland. <i>Molecular Biology Reports</i> , 2020, 47, 3929-3935.	2.3	6
13	Evaluation of PCR markers for <i>Phytophthora infestans</i> mating type determination. <i>European Journal of Plant Pathology</i> , 2018, 152, 33-44.	1.7	8
14	QTL for tuber morphology traits in diploid potato. <i>Journal of Applied Genetics</i> , 2018, 59, 123-132.	1.9	24
15	Quantitative trait loci for tuber blackspot bruise and enzymatic discoloration susceptibility in diploid potato. <i>Molecular Genetics and Genomics</i> , 2018, 293, 331-342.	2.1	12
16	Diversity of <i>AvrEcnt1</i> and <i>AvrSmira1</i> effector genes in Polish and Norwegian populations of <i>Phytophthora infestans</i> . <i>Plant Pathology</i> , 2018, 67, 1792-1802.	2.4	9
17	Novel gene <i>Sen2</i> conferring broad-spectrum resistance to <i>Synchytrium endobioticum</i> mapped to potato chromosome XI. <i>Theoretical and Applied Genetics</i> , 2018, 131, 2321-2331.	3.6	22
18	Characterization of <i>Dickeya</i> and <i>Pectobacterium</i> strains obtained from diseased potato plants in different climatic conditions of Norway and Poland. <i>European Journal of Plant Pathology</i> , 2017, 148, 839-851.	1.7	42

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19	Expression of the Potato Late Blight Resistance Gene <i>Rpi-phu1</i> and <i>Phytophthora infestans</i> Effectors in the Compatible and Incompatible Interactions in Potato. <i>Phytopathology</i> , 2017, 107, 740-748.	2.2	25
20	BC1 and F1 Progeny from <i>Solanum</i> <i>microocanum</i> (+) <i>S. tuberosum</i> Somatic Hybrids, Autofused 4 \times <i>S. microocanum</i> and Cultivated Potato. <i>American Journal of Potato Research</i> , 2017, 94, 323-333.	0.9	7
21	Quantitative trait loci affecting intensity of violet flower colour in potato. <i>Euphytica</i> , 2017, 213, 1.	1.2	2
22	<i>Phytophthora infestans</i> : Isolation of Pure Cultures, Storage and Inoculum Preparation. <i>Plant Breeding and Seed Science</i> , 2017, 76, 9-15.	0.1	7
23	Laboratory Assessment of Potato Resistance to <i>Phytophthora infestans</i> . <i>Plant Breeding and Seed Science</i> , 2017, 76, 17-23.	0.1	8
24	The effect of drought stress on the leaf relative water content and tuber yield of a half-sib family of <i>Katahdin</i> TM -derived potato cultivars. <i>Breeding Science</i> , 2016, 66, 328-331.	1.9	149
25	Diversity of <i>Fusarium</i> spp. associated with dry rot of potato tubers in Poland. <i>European Journal of Plant Pathology</i> , 2016, 145, 871-884.	1.7	59
26	Genetic composition of interspecific potato somatic hybrids and autofused 4 \times plants evaluated by DArT and cytoplasmic DNA markers. <i>Plant Cell Reports</i> , 2016, 35, 1345-1358.	5.6	29
27	Potato cultivation system affects population structure of <i>Phytophthora infestans</i> . <i>Fungal Ecology</i> , 2016, 20, 132-143.	1.6	22
28	Mapping of quantitative trait loci for tuber starch and leaf sucrose contents in diploid potato. <i>Theoretical and Applied Genetics</i> , 2016, 129, 131-140.	3.6	26
29	Fine mapping of the <i>Rpi-rzc1</i> gene conferring broad-spectrum resistance to potato late blight. <i>European Journal of Plant Pathology</i> , 2015, 143, 193-198.	1.7	14
30	Novel candidate genes <i>AuxRP</i> and <i>Hsp90</i> influence the chip color of potato tubers. <i>Molecular Breeding</i> , 2015, 35, 224.	2.1	28
31	R2-like Gene Contributes to Resistance to <i>Phytophthora infestans</i> in Polish Potato Cultivar Bzura. <i>American Journal of Potato Research</i> , 2015, 92, 350-358.	0.9	23
32	Diversity of <i>Phytophthora infestans</i> from Poland. <i>Plant Pathology</i> , 2014, 63, 203-211.	2.4	38
33	Hypersensitive response to Potato virus Y in potato cultivar <i>Smira</i> is conferred by the <i>Ny-Smira</i> gene located on the long arm of chromosome IX. <i>Molecular Breeding</i> , 2014, 34, 471-480.	2.1	24
34	A locus conferring effective late blight resistance in potato cultivar <i>Smira</i> maps to chromosome XI. <i>Theoretical and Applied Genetics</i> , 2014, 127, 647-657.	3.6	28
35	Resistance to <i>Phytophthora infestans</i> in Three <i>Solanum nigrum</i> F3 Families. <i>Plant Breeding and Seed Science</i> , 2014, 66, 63-73.	0.1	4
36	Resistance gene enrichment sequencing (RESeq) enables reannotation of the <i>LRR</i> gene family from sequenced plant genomes and rapid mapping of resistance loci in segregating populations. <i>Plant Journal</i> , 2013, 76, 530-544.	5.7	367

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37	Interspecific somatic hybrids <i>Solanum villosum</i> (+) <i>S. tuberosum</i> , resistant to <i>Phytophthora infestans</i> . <i>Journal of Plant Physiology</i> , 2013, 170, 1541-1548.	3.5	20
38	Development of somatic hybrids <i>Solanum</i> – <i>Michoacanum</i> Bitter. (Rydb.) (+) <i>S. tuberosum</i> L. and autofused 4x <i>S.</i> – <i>Michoacanum</i> plants as potential sources of late blight resistance for potato breeding. <i>Plant Cell Reports</i> , 2013, 32, 1231-1241.	5.6	27
39	Influence of genetic background and plant age on expression of the potato late blight resistance gene <i>Rpi-phu1</i> during incompatible interactions with <i>Phytophthora infestans</i> . <i>Plant Pathology</i> , 2013, 62, 1072-1080.	2.4	20
40	The influence of long-term storage in liquid nitrogen on survival and pathogenicity of <i>Phytophthora infestans</i> isolates. <i>Journal of Plant Protection Research</i> , 2012, 52, 479-485.	1.0	3
41	Late blight resistance gene from <i>Solanum ruiz-ceballosii</i> is located on potato chromosome X and linked to violet flower colour. <i>BMC Genetics</i> , 2012, 13, 11.	2.7	44
42	A resistance gene against potato late blight originating from <i>Solanum</i> – <i>Michoacanum</i> maps to potato chromosome VII. <i>Theoretical and Applied Genetics</i> , 2012, 124, 397-406.	3.6	46
43	Marker-assisted selection of diploid and tetraploid potatoes carrying <i>Rpi-phu1</i> , a major gene for resistance to <i>Phytophthora infestans</i> . <i>Journal of Applied Genetics</i> , 2010, 51, 133-140.	1.9	25
44	<i>Rpi-vnt1.1</i> , a <i>Tm-2</i> Homolog from <i>Solanum venturii</i> , Confers Resistance to Potato Late Blight. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 589-600.	2.6	194
45	Recognition of <i>Phytophthora infestans</i> Avr4 by potato R4 is triggered by C-terminal domains comprising W motifs. <i>Molecular Plant Pathology</i> , 2009, 10, 611-620.	4.2	22
46	Tagging quantitative trait loci for dormancy, tuber shape, regularity of tuber shape, eye depth and flesh colour in diploid potato originated from six <i>Solanum</i> species. <i>Plant Breeding</i> , 2008, 127, 49-55.	1.9	32
47	Tagging QTLs for late blight resistance and plant maturity from diploid wild relatives in a cultivated potato (<i>Solanum tuberosum</i>) background. <i>Theoretical and Applied Genetics</i> , 2007, 115, 101-112.	3.6	32
48	Mating Type, Virulence, Aggressiveness and Metalaxyl Resistance of Isolates of <i>Phytophthora infestans</i> in Poland. <i>Potato Research</i> , 2007, 49, 155-166.	2.7	24
49	The novel, major locus <i>Rpi-phu1</i> for late blight resistance maps to potato chromosome IX and is not correlated with long vegetation period. <i>Theoretical and Applied Genetics</i> , 2006, 113, 685-695.	3.6	95
50	Genetic factors encoding resistance to late blight caused by <i>Phytophthora infestans</i> (Mont.) de Bary on the potato genetic map. <i>Cellular and Molecular Biology Letters</i> , 2004, 9, 855-67.	7.0	4